

Resource Summary Report

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PDBe - Protein Data Bank in Europe

RRID:SCR_004312

Type: Tool

Proper Citation

PDBe - Protein Data Bank in Europe (RRID:SCR_004312)

Resource Information

URL: <http://www.ebi.ac.uk/pdbe/>

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Description: The European resource for the collection, organization and dissemination of data on biological macromolecular structures. In collaboration with the other worldwide Protein Data Bank (wwPDB) partners - the Research Collaboratory for Structural Bioinformatics (RCSB) and BioMagResBank (BMRB) in the USA and the Protein Data Bank of Japan (PDBj) - they work to collate, maintain and provide access to the global repository of macromolecular structure data. The main objectives of the work at PDBe are: * to provide an integrated resource of high-quality macromolecular structures and related data and make it available to the biomedical community via intuitive user interfaces. * to maintain in-house expertise in all the major structure-determination techniques (X-ray, NMR and EM) in order to stay abreast of technical and methodological developments in these fields, and to work with the community on issues of mutual interest (such as data representation, harvesting, formats and standards, or validation of structural data). * to provide high-quality deposition and annotation facilities for structural data as one of the wwPDB deposition sites. Several sophisticated tools are also available for the structural analysis of macromolecules.

Abbreviations: PDBe

Synonyms: Protein DataBank Europe, Protein DataBank in Europe, PDBe, Protein Data Bank in Europe, Protein Data Bank Europe, Macromolecular Structure Database

Resource Type: data or information resource, database, service resource, storage service resource, data repository

Defining Citation: [PMID:21045060](#), [PMID:21460450](#), [PMID:19858099](#)

Keywords: x-ray, nmr, cryo-em, hybrid method, dna, protein, rna, sugar, ligand, virus, compound, fold, enzyme, 3d spatial image, structure, macromolecule, protein-protein interaction, gold standard, bio.tools

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CCP4

Resource Name: PDBe - Protein Data Bank in Europe

Resource ID: SCR_004312

Alternate IDs: r3d100012791, biotools:pdbe, nlx_32372

Alternate URLs: <https://bio.tools/pdbe>, <https://doi.org/10.17616/R3J226>

Record Creation Time: 20220129T080223+0000

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Ratings and Alerts

No rating or validation information has been found for PDBe - Protein Data Bank in Europe.

No alerts have been found for PDBe - Protein Data Bank in Europe.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 55 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Querino Lima Afonso M, et al. (2026) PDBe: enhanced structural data exploration to facilitate discovery. Nucleic acids research, 54(D1), D440.

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Dodonova SO, et al. (2019) Structure of the Ty3/Gypsy retrotransposon capsid and the evolution of retroviruses. *Proceedings of the National Academy of Sciences of the United States of America*, 116(20), 10048.

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Cook CE, et al. (2019) The European Bioinformatics Institute in 2018: tools, infrastructure and training. *Nucleic acids research*, 47(D1), D15.

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Kolesnikova O, et al. (2018) Molecular structure of promoter-bound yeast TFIID. *Nature communications*, 9(1), 4666.

De Robertis M, et al. (2018) In Vivo Evaluation of a New Recombinant Hyaluronidase to Improve Gene Electro-Transfer Protocols for DNA-Based Drug Delivery against Cancer. *Cancers*, 10(11).